

| | | |
|------------|--------------------------------------------------------------|-----|
| z219c-a_-r | MRVSGVLRLLALIFAIVTTWMFIRSYMS-FSMKTIRLPRWLASP---- <td>45</td> | 45 |
| MMU72677_1 | ----- | |
| 219_HUMAN | MRLAGPLRIVVLVVSQVGTWIVVSILLGGPGSGFPRIQQLFTSPESSVTA | 50 |
| D87120_1_ | MRVAGAAKLVVAVAVFLLTFYVISQVFE- IKMDASLGNLFARSALDTAAR | 49 |
| z219c-a_-r | EIQVKKYKCGLIKPCPANYFAFKICSGAANVVGPTMCFEDRMIMSPVKNN | 95 |
| MMU72677_1 | ----- | |
| 219_HUMAN | APRARKYKCGLPQPCPEEHLAFRVVSGAANVIGPKICLEDKMLMSSVKDN | 100 |
| D87120_1_ | STKPPRYKCGISKACPEKHFAFKMASGAANVVGPKICLEDNVLMMSGVKNN | 99 |
| z219c-a_-r | VGRGLNIALVNGTTGAVLGQKAFDMYSGDVMHLVKFLKEIPGGALVLVAS | 145 |
| MMU72677_1 | -----TGQVMKKDSFDMYSGDPQLLLNFLTEIPDSTLVLVAS | 37 |
| 219_HUMAN | VGRGLNIALVNGVSGELIEARAFDMWAGDVNDLLKFIRPLHEGTLVFVAS | 150 |
| D87120_1_ | VGRGINVALANGKTGEVLDTKYFDMWGGDVAPFIEFLKAIQDGTIVLMGT | 149 |
| z219c-a_-r | YDDPGTKMNDESRLKFSDLGSSYAKQLGFRDSWVFIGAKDLRGKSPFEQF | 195 |
| MMU72677_1 | YDDPGTKMNDKIKTLFSNLGSSYAKQLGFRDSWVFIGAKDLKSKSPYEQF | 87 |
| 219_HUMAN | YDDPATKMNEETRKLFSSELGSRNAKELAFRDSWVFIGAKGVQNKSPFEQH | 200 |
| D87120_1_ | YDDGATKLNDEARRLIADLGSTSITNLGFRDNWVFCGGKGIKTKSPFEQH | 199 |
| z219c-a_-r | LKNSPDTNKYEGWPELLEMEGCMPPKPF-- | 223 |
| MMU72677_1 | LKNNPETNKYDGWPELLELEGCVPRKVM-- | 115 |
| 219_HUMAN | VKNSKHSNKYEGCPEALEMEGCIPRRSTAS | 230 |
| D87120_1_ | IKNNKDTNKYEGWPEVVEMEGCIPQKQD-- | 227 |

Figure